# Analysis Facilities for the HL-LHC

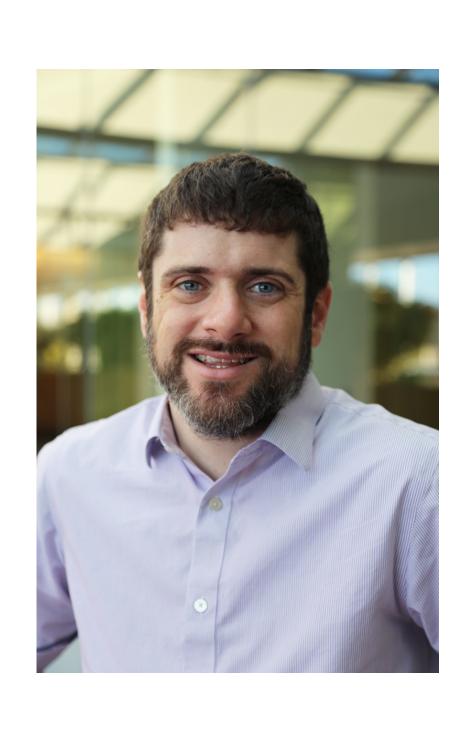






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#### A bit about me



I have been working on CMS computing since 2006 and the OSG since 2008.

- Helped introduce data federations based on XRootD to the community as part of the AAA project.
- Accordingly, this presentation is going to be strongly colored by my experience (e.g., LHC-centric, CMS-centric).

Currently at the Morgridge Institute for Research where I work closely with Miron Livny and the Center for High Throughput Computing.

• PhD is in Math and CS – will not be in danger of covering any physics topics in this talk!

# What is an Analysis Facility, anyway?

When I say 'analysis facility,' what do I mean?

- People, software / services, and hardware meant to support analysis activities for an experiment aggregated into a coherent whole.
- Services includes:
  - Access to experimental data products.
  - Storage space for per-group or per-user data (often ntuples).
  - Access to significant computing resources.
- I promised to not talk about physics software, but obviously this is ROOT and the growing Python-based ecosystem.
- Computing hardware (currently) looks like most of our computing facilities: CPUs and disks. More emphasis on high-data-rates. However, there's a growing need for GPUs.



# Great Example - USCMS LPC

What exists today at the USCMS LPC?

- Dedicated staff to help.
- Users can login to hosts via SSH and do modest technical work.
  - Any large-scale computation is done via HTCondor.
- Filesystems for input and output currently using EOS.
  - Uses the same authentication mechanism as CMS - X.509(GSI/VOMS).

# LPC Services

Compute Interface	SSH host & Batch System
Batch System	HTCondor
Data Interface	POSIX file-like
Input products	EOS disk / offsite XRootD
Data Storage	EOS disk



# Why are we interested in Analysis Facilities?

The LHC has been running for years - why are we interested in an AF now?

- HL-LHC will have at least one order-magnitude increase in event count, if not two.
  - Premise: Work previously done on the laptop will now require significant computing resources.
- We want analysts to use significant resources interactively, similarly to how they use their laptops today.
  - Batch jobs even when they startup quickly have a distinct experience from interactive use.
- We want new <u>services</u> to be available to users as well as new <u>resource types</u>.



# **Columnar Analysis**

We also want to explore what services are needed in order to support columnar analysis.

- Operations should be expressed on arrays of data, not individual events.
- Represent HEP data as arrays in memory, allowing us to leverage existing libraries that work on vector data (think: numpy).
- Track non-uniformity ("jaggedness") of arrays in a separate array.
- No C++-style objects in memory!

# Run Physics Analysis on a GPU

Notebook from Joosep Pata's tutorial at PyHEP 2020



See also: https://coffeateam.github.io/coffea/



# Our Work

#### The Nebraska CMS-AF

We are working to provide a new facility at the CMS Tier-2 site at Nebraska. Goal:

- Any CMS user can start a JupyterLab interface and get computational resources through the familiar "Dask" interface.
- Have all auth $\{n,z\}$  be based on web single-sign-on – not X.509 client certificates.
- As the community grows, add new services to the stack.

# **CMS-AF Services**

Compute Interface	Jupyter Notebooks & Dask Scheduler
Task Scheduler	Dask
Data Interface	Object-store like
Input products	CMS AAA through XCache
Data Storage	??? Unresolved.

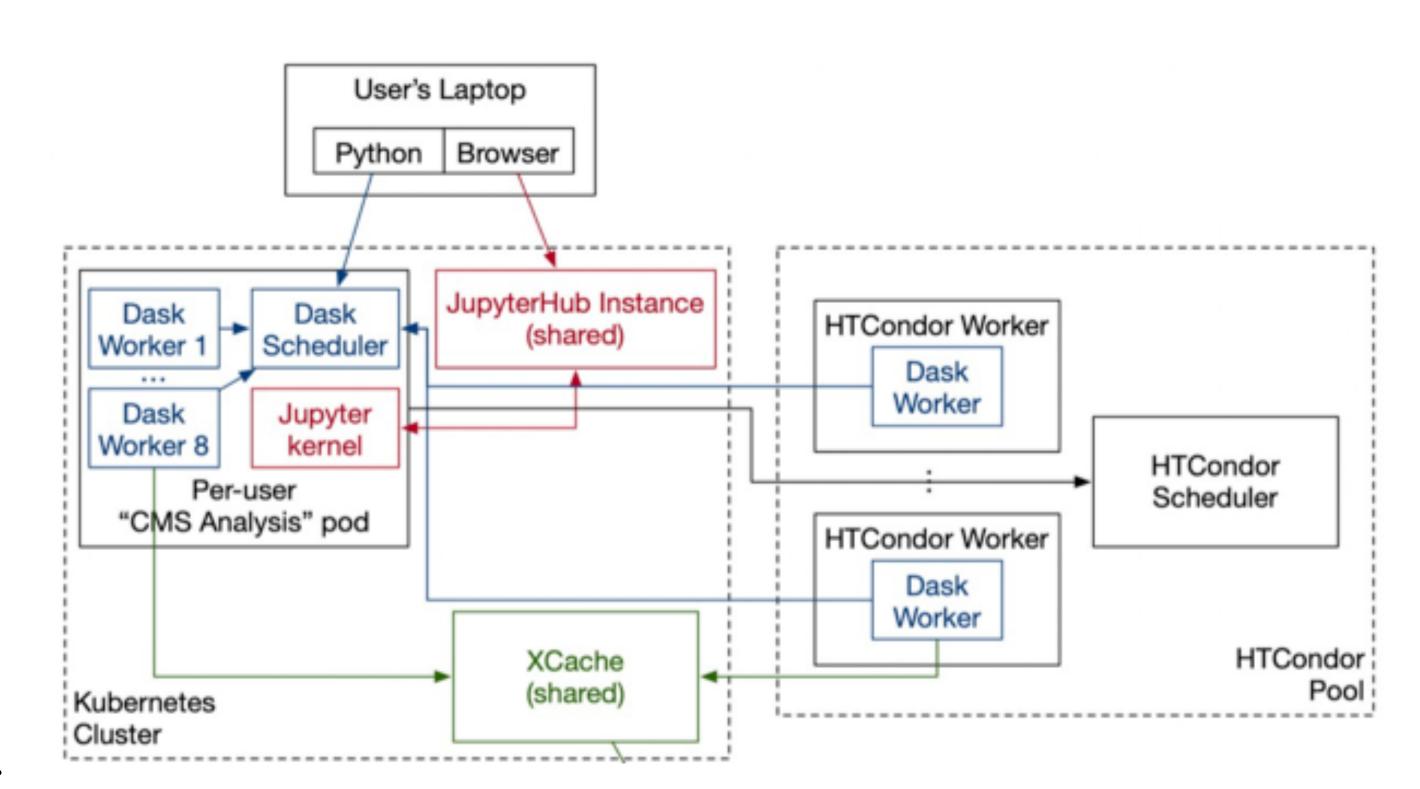
YouTube demo.



### First service – the JupyterLab & Dask Scheduler

Our first service – functioning now, but aiming to scale to all of USCMS – is a Dask scheduler.

- Completely focused on interactive use cases. Dask workers are launched as soon as the user's notebook is available.
  - "A chicken in every pot and 8 cores for every analyst".
- Beyond the resources started in Kubernetes, we launch additional workers in HTCondor.
  - Resources beyond the 'standard allocation' are allocated by HTCondor via the standard fairshare mechanism.
- We feel it is essential that these analysis resources coexist with the existing resources – we can't afford partitioning a standalone facility.

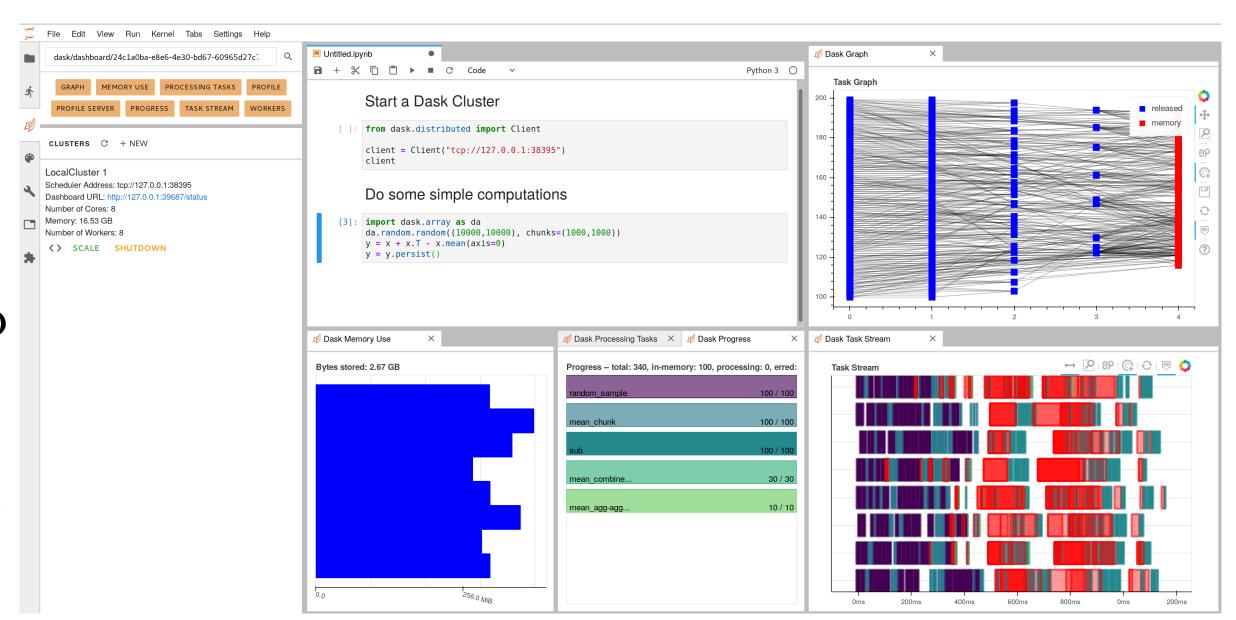


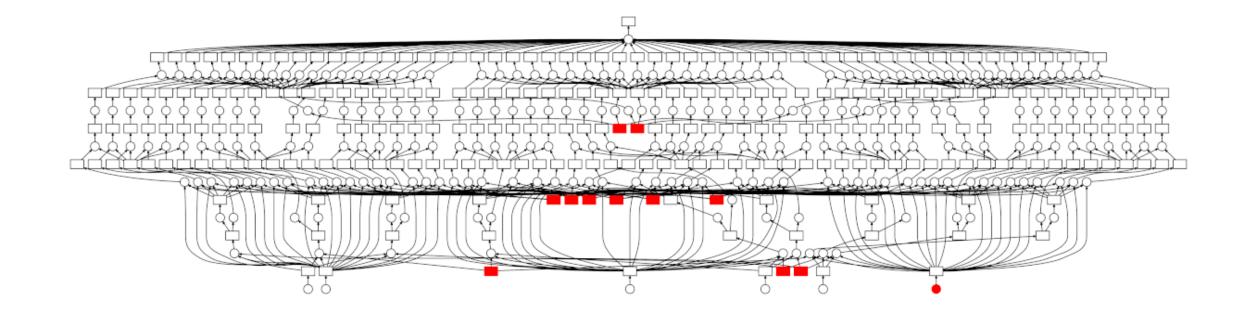
#### An Aside on Dask

Dask provides a simple task-based parallelism interface in Python.

- Tasks are scheduled out in milliseconds to dedicated worker processes.
- Dask workers keep intermediate results in memory, providing for data locality and minimizing data movement.

Dask provides interactive use, which is perhaps the biggest difference compared to today's batch-system-based solution.





## **Beyond Dask**

What are the minimal services beyond Dask?

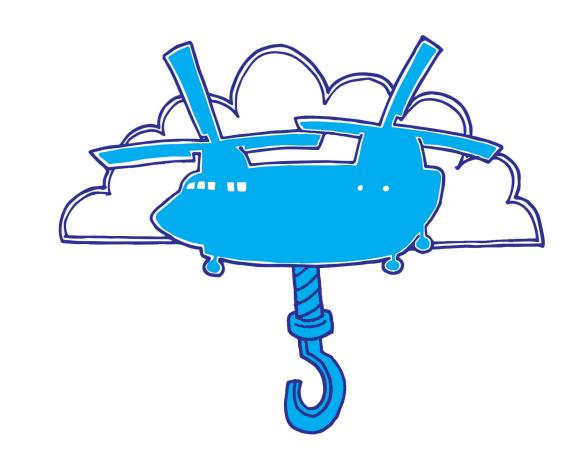
- Authentication and authorization based on OAuth2 ensure that only CMS users can access the AF.
- Data is delivered through XCache: when tasks are started, they are started with credentials that can read from the onsite XCache, which proxy out to the wider CMS data federation.
- Local storage for code / notebooks.

Big open question: what should be done about user data?

# Next up: data handling

We think of the Dask scheduler as the initial service for the CMS-AF. What's next?

- Derivation of columns & ntuples: Most analyses can't use experiment-produced ntuples directly but need to generate them from an analysis format. We are looking at ServiceX, developed by IRIS-HEP. ServiceX derives columns on demand from external datasets and can deliver them to object storage.
- Actual column store: Right now, all events are still kept in a file structure. We are looking at using SkyHook DM to store ntuples in a dataset as a "table-like" structure -- but add database-like primitives (SELECT/PROJECT/FILTER).
  - Particularly, being able to augment existing tables (datasets) with additional columns is seen as a killer feature.



#### Take-Home

What's the message for the larger Snowmass community?

- Analysis facilities have been with us for awhile they have been important and will be important going forward.
  - Current facilities are largely based on the "batch system + filesystem" paradigm.
- We don't have to stand still! There are a variety of other approaches (interactive notebooks, task-based, column stores) to explore.
  - Nebraska is working toward a specific vision, developing initially a Dask service.
  - There are several other parallel efforts I'm aware of (CERN's SWAN, Vanderbilt, FNAL Elastic Compute Facility).

# So, how should we all proceed?



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